

QY 121 TSETSDIQEYGRVRAAAGSAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
 Db 121 TSETSDIQEYGRVRAAAGSAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
 QY 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
 Db 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
 QY 241 ABIYQMLDRRSORSEERCVEIP 263
 Db 241 ABIYQMLDRRSORSEERCVEIP 263

RESULT 2

US-10-293-832-28
 ; Sequence 28, Application US/10293832
 ; Publication No. US20030180752A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Liu, Wei
 ; APPLICANT: Fouser, Lynette
 ; APPLICANT: Spaulding, Vikki
 ; TITLE OF INVENTION: TYPE 2 CYTOKINE RECEPTOR AND NUCLEIC ACIDS ENCODING SAME
 ; FILE REFERENCE: 22058-546
 ; CURRENT APPLICATION NUMBER: US/10/293,832
 ; PRIOR FILING DATE: 2002-11-12
 ; PRIOR APPLICATION NUMBER: US 60/332,366
 ; PRIOR FILING DATE: 2001-11-09
 ; NUMBER OF SEQ ID NOS: 33
 ; SOFTWARE: PatentIn version 3.2
 ; SEQ ID NO 28
 ; LENGTH: 263
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-10-293-832-28

Query Match 100.0%; Score 1432; DB 12; Length 263;
 Best Local Similarity 100.0%; Pred. No. 4e-138;
 Matches 263; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MPMKHCFLGLISFFLTGAGTOSTHESLKQVQFQSRNFHILQWQGRALTNSSVY 60
 Db 1 MPMKHCFLGLISFFLTGAGTOSTHESLKQVQFQSRNFHILQWQGRALTNSSVY 60
 QY 61 FVOYKIMFSCMSKSHQKSGCWHISCNPPGCRITLAKYQKQWKNKEDCWGTQELSCDL 120
 Db 61 FVOYKIMFSCMSKSHQKSGCWHISCNPPGCRITLAKYQKQWKNKEDCWGTQELSCDL 120
 QY 121 TSETSDIQEYGRVRAAAGSAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
 Db 121 TSETSDIQEYGRVRAAAGSAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
 QY 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
 Db 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
 QY 241 ABIYQMLDRRSORSEERCVEIP 263
 Db 241 ABIYQMLDRRSORSEERCVEIP 263

RESULT 3

US-10-312-088-42
 ; Sequence 42, Application US/10312088
 ; Publication No. US20030219862A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Agarwal, Pankaj
 ; APPLICANT: Cogswell, John P.
 ; APPLICANT: Kabnic, Karen S.
 ; APPLICANT: Lai, Ying-Ta
 ; APPLICANT: Martensen, Shelby A.
 ; APPLICANT: Murdock, Paul R.
 ; APPLICANT: Smith, Randall F.

; APPLICANT: Strum, Jay C.
 ; APPLICANT: Xiang, Zhaoying
 ; APPLICANT: Xie, Qing
 ; APPLICANT: Rizni, Safia K.
 ; TITLE OF INVENTION: NOVEL COMPOUNDS
 ; FILE REFERENCE: GP50029
 ; CURRENT APPLICATION NUMBER: US/10/312,088
 ; CURRENT FILING DATE: 2002-12-20
 ; PRIOR APPLICATION NUMBER: PCT/US01/19929
 ; PRIOR FILING DATE: 2001-06-22
 ; PRIOR APPLICATION NUMBER: 60/213,161
 ; PRIOR FILING DATE: 2000-06-22
 ; PRIOR APPLICATION NUMBER: 60/213,156
 ; PRIOR FILING DATE: 2000-06-22
 ; NUMBER OF SEQ ID NOS: 44
 ; SOFTWARE: Fast-Seq for Windows Version 4.0
 ; SEQ ID NO 42
 ; LENGTH: 263
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-10-312-088-42

Query Match 100.0%; Score 1432; DB 12; Length 263;
 Best Local Similarity 100.0%; Pred. No. 4e-138;
 Matches 263; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MPMKHCFLGLISFFLTGAGTOSTHESLKQVQFQSRNFHILQWQGRALTNSSVY 60
 Db 1 MPMKHCFLGLISFFLTGAGTOSTHESLKQVQFQSRNFHILQWQGRALTNSSVY 60
 QY 61 FVOYKIMFSCMSKSHQKSGCWHISCNPPGCRITLAKYQKQWKNKEDCWGTQELSCDL 120
 Db 61 FVOYKIMFSCMSKSHQKSGCWHISCNPPGCRITLAKYQKQWKNKEDCWGTQELSCDL 120
 QY 121 TSETSDIQEYGRVRAAAGSAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
 Db 121 TSETSDIQEYGRVRAAAGSAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
 QY 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
 Db 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
 QY 241 ABIYQMLDRRSORSEERCVEIP 263
 Db 241 ABIYQMLDRRSORSEERCVEIP 263

RESULT 4

US-09-961-404-6
 ; Sequence 6, Application US/09961404
 ; Publication No. US2003022827A1
 ; GENERAL INFORMATION:
 ; APPLICANT: WEISS, BERTRAM
 ; APPLICANT: SABAT, ROBERT
 ; APPLICANT: ASADULLAH, KHUSRU
 ; APPLICANT: TOSCHI, LUISELLA
 ; TITLE OF INVENTION: THREE NEW MEMBERS OF THE CYTOKINE RECEPTOR
 ; FILE REFERENCE: SCH-1788
 ; CURRENT APPLICATION NUMBER: US/09/961,404
 ; CURRENT FILING DATE: 2001-09-25
 ; NUMBER OF SEQ ID NOS: 19
 ; SOFTWARE: PatentIn Ver. 2.1
 ; SEQ ID NO 6
 ; LENGTH: 263
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-09-961-404-6

Query Match 99.7%; Score 1428; DB 11; Length 263;
 Best Local Similarity 99.6%; Pred. No. 1e-137;
 Matches 262; Conservative 1; Mismatches 0; Indels 0; Gaps 0;